FOS course

Molecular Data Science



In person teaching



Key to education but now a fragile luxury.

Measures



- Stay home and test if you have symptoms
- Use face mask in building outside class
- Keep distance
- Adhere to rules outside the course
- Vaccination at LUMC without appointment
- Hybrid teaching in case of symptoms
- How is tbd? Presentations: central in Teams, Practicals: with buddy in Teams?
- Be flexible & mild for yourself and each other

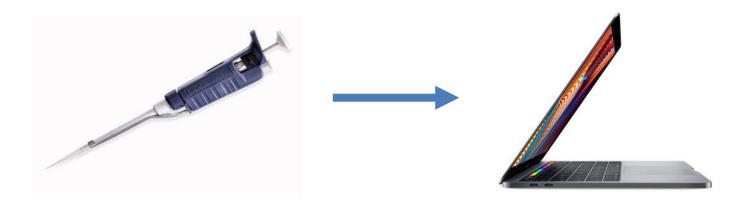






Molecular Data Science

Solving a biological puzzles without touching a pipette



Data drive the modern life sciences

"Tomorrow's discoveries will be made by today's early-career researchers. ... We also hope to be guided by them, so that we can meet their needs when it comes to publishing their work, as research becomes more datarich and computationally heavy."



Magdalena Skipper, Editor-in-Chief Nature

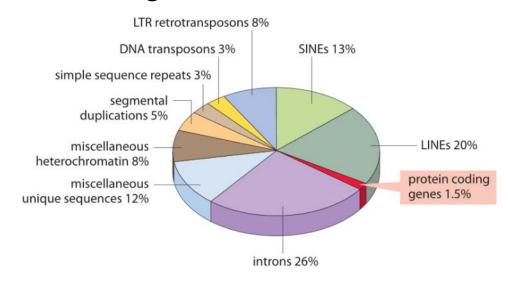


Many discoveries remain

The human genome

- ~ 3.2 billion bases (billion = miljard)
- ~ 22 thousand protein coding genes
- ~ 98% of human genome is non-coding

... and we hardly have a clue.



Many discoveries remain

Our human DNA is uncharted territory

- Dedicated paper on 5400 genes.
- 90% of papers just on 2,000 genes.
- 2015: 50% of research devoted on 3,000 genes known by 1991



Data intensive

Acknowledge lack of knowledge: genome-wide



Gene
PCR
Chosen by researcher



Genome-wide Array All current knowledge

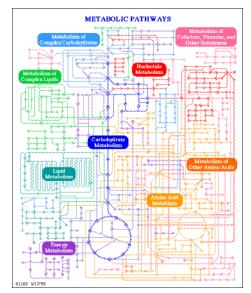


Whole genome
Sequencing
Everything there is



Data intensive

- Acknowledge lack of knowledge: genome-wide
- From reductionist (1 gene) to the system (all genes)



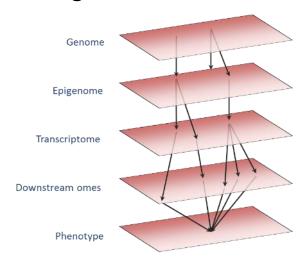
Acknowledges

- Complexity of biology
- Advantage of whole picture



Data intensive

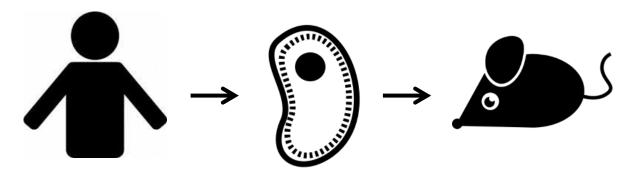
- Acknowledge lack of knowledge: genome-wide
- From reductionist (1 gene) to the system (all genes)
- Integrative: combine levels to trace processes





Data intensive

- Acknowledge lack of knowledge: genome-wide
- From reductionist (1 gene) to the system (all genes)
- Integrative: combine levels to trace processes
- Using natural variation (instead of experimental) in large-scale population studies





Who

Tutors

- Ingrid Meulenbelt & Bas Heijmans (coordinators)
- 15 others

You

- 14 Master students Biomedical Sciences
- 2 PhD students, 1 postdoc, 1 data science teacher (first 2.5 weeks)



The course

- Learning by doing: analyzing all kind of omics data → 12 days
- Applying knowledge: develop molecular data science project → 8 days



Tools

RStudio Cloud



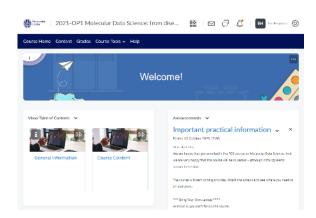
Data science without the hardware hassles

Discord, Teams...





Brightspace





Learning objectives



Knowledge and understanding

- Knows how large-scale molecular data can inform on mechanisms and risk of common diseases.
- Has insight in modern data analysis methods used to discover molecular signatures of disease phenotypes in genetic, epigenetic, gene expression, and metabolomics data sets.



Applying knowledge and understanding

- Get hands-on experience in the analysis and interpretation of genetic, epigenetic, gene expression, and metabolomics data sets.
- Shows the ability to develop new researcher project in the field of ageing using molecular data science including background, hypothesis, pilot data, objectives, study design, work plan, and expected outcomes (e.g. causality).
- Can perform analyses to generate pilot data in order to critically appraise and, if necessary, reformulate a hypothesis.



Communication

- Shows communication skills to clearly and convincingly present and defend a research proposal.
- Is able to respond constructively to questions/feedback and connecting this
 feedback to his/her own position regarding his/her own research and in
 doing so showing an open, self-critical yet firm and self-confident attitude.



Learning skills

- Shows professional conduct: being critical yet constructive and eager to improve oneself and in doing so contributes to the learning process of the other students.
- Critically and constructively discusses research proposals of peers.



Pay close attention Assessment



Assessment

- 1. Handing in assignments. Individually assessed (0%, P/F).
- 2. Contribute to interim evaluation of student participation and development during workgroups (0%).
- 3. Fill out project proposal form (for reflective assignment) (0%, P/F).
- 4. Presentation project proposal (background, hypothesis, pilot data, objectives, study design, workplan, expected outcomes) (45%).
- 5. Active and critical participation during discussion after project presentations of peers (15%).
- 6. Reflective assignment that shows mastering key aspects of development of research proposal in molecular data science and addressing points raised during peer review (40%).

Evaluation

Who will it be?

(2 are needed)





Part of the Master Track Data-driven Research

